

# Sequence Listing

<110> Goddard, Audrey  
 Godowski, Paul J.  
 Gurney, Austin L.  
 Watanabe, Colin K.  
 Wood, William I.

<120> NOVEL POLYPEPTIDES HAVING SEQUENCE SIMILARITY TO  
 CYTOKINE RECEPTORS AND NUCLEIC ACIDS ENCODING THE SAME

<130> P3121R1

<140> US 09/964,994

<141> 2001-09-26

<150> PCT/US00/08439

<151> 2000-03-30

<150> PCT/US01/06520

<151> 2001-02-28

<150> US 60/191,015

<151> 2000-03-21

<150> US 09/941,992

<151> 2001-08-28

<160> 7

<210> 1

<211> 1318

<212> DNA

<213> Homo Sapien

<400> 1

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 35 40 45  
 Trp Gln Pro Gly Arg Ala Leu Thr Gly Asn Ser Ser Val Tyr Phe  
 50 55 60  
 Val Gln Tyr Lys Ile Met Phe Ser Cys Ser Met Lys Ser Ser His  
 65 70 75  
 Gln Lys Pro Ser Gly Cys Trp Gln His Ile Ser Cys Asn Phe Pro  
 80 85 90  
 Gly Cys Arg Thr Leu Ala Lys Tyr Gly Gln Arg Gln Trp Lys Asn  
 95 100 105  
 Lys Glu Asp Cys Trp Gly Thr Gln Glu Leu Ser Cys Asp Leu Thr  
 110 115 120  
 Ser Glu Thr Ser Asp Ile Gln Glu Pro Tyr Tyr Gly Arg Val Arg  
 125 130 135  
 Ala Ala Ser Ala Gly Ser Tyr Ser Glu Trp Ser Met Thr Pro Arg  
 140 145 150  
 Phe Thr Pro Trp Trp Glu Thr Lys Ile Asp Pro Pro Val Met Asn  
 155 160 165

Ile Thr Gln Val Asn Gly Ser Leu Leu Val Ile Leu His Ala Pro  
 170 175 180  
 Asn Leu Pro Tyr Arg Tyr Gln Lys Glu Lys Asn Val Ser Ile Glu  
 185 190 195  
 Asp Tyr Tyr Glu Leu Leu Tyr Arg Val Phe Ile Ile Asn Asn Ser  
 200 205 210  
 Leu Glu Lys Glu Gln Lys Val Tyr Glu Gly Ala His Arg Ala Val  
 215 220 225  
 Glu Ile Glu Ala Leu Thr Pro His Ser Ser Tyr Cys Val Val Ala  
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20 25 30

Val Val Leu Pro Cys Leu Gly Phe Thr Leu Leu Leu Trp Ser Gln  
35 40 45

Val Ser Gly Ala Gln Gly Gln Glu Phe His Phe Gly Pro Cys Gln  
50 55 60

Val Lys Gly Val Val Pro Gln Lys Leu Trp Glu Ala Phe Trp Ala  
65 70 75

Val Lys Asp Thr Met Gln Ala Gln Asp Asn Ile Thr Ser Ala Arg  
80 85 90

Leu Leu Gln Gln Glu Val Leu Gln Asn Val Ser Asp Ala Glu Ser  
95 100 105

Cys Tyr Leu Val His Thr Leu Leu Glu Phe Tyr Leu Lys Thr Val  
110 115 120

Phe Lys Asn His His Asn Arg Thr Val Glu Val Arg Thr Leu Lys  
125 . 130 135

Ser Phe Ser Thr Leu Ala Asn Asn Phe Val Leu Ile Val Ser Gln  
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Leu Gln Pro Ser Gln Glu Asn Glu Met Phe Ser Ile Arg Asp Ser  
155 160 165

Ala His Arg Arg Phe Leu Leu Phe Arg Arg Ala Phe Lys Gln Leu  
170 175 180

Asp Val Glu Ala Ala Leu Thr Lys Ala Leu Gly Glu Val Asp Ile  
185 190 195

Leu Leu Thr Trp Met Gln Lys Phe Tyr Lys Leu  
200 205